

<!--StartFragment-->RESULT 5
 BH789726/c

LOCUS BH789726 65 bp DNA linear GSS 02-APR-2002

DEFINITION SALK_044532.19.55.x Arabidopsis thaliana TDNA insertion lines
 Arabidopsis thaliana genomic clone SALK_044532.19.55.x, genomic survey sequence.

ACCESSION BH789726

VERSION BH789726.1 GI:19882824

KEYWORDS GSS.

SOURCE Arabidopsis thaliana (thale cress)

ORGANISM Arabidopsis thaliana
 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
 Spermatophyta; Magnoliophyta; eudicots; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

REFERENCE 1 (bases 1 to 65)

AUTHORS Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R.,
 Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L.,
 Shinn,P., Zimmerman,J. and Ecker,J.R.

TITLE A Sequence-Indexed Library of Insertion Mutations in the
 Arabidopsis Genome

JOURNAL Unpublished (2001)

COMMENT Contact: Joseph R. Ecker
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 The Salk Institute for Biological Studies
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 This is single pass sequence recovered from the left border of
 TDNA. This sequence lies within an annotated exon of At3g41627.
 Class: TDNA tagged.

FEATURES Location/Qualifiers

source 1..65
 /organism="Arabidopsis thaliana"
 /mol_type="genomic DNA"
 /ecotype="Col-0"
 /db_xref="taxon:3702"
 /clone="SALK_044532.19.55.x"
 /clone_lib="Arabidopsis thaliana TDNA insertion lines"
 /note="PCR was performed on Arabidopsis thaliana lines
 each of which contains one or more TDNA insertion
 elements. The resultant fragment for each line was
 directly sequenced to determine the genomic sequence at
 the site of insertion. Details of the protocols used can
 be found at http://signal.salk.edu/tdna_protocols.html"

ORIGIN

Query Match	100.0%	Score	12	DB	15	Length	65
Best Local Similarity	100.0%	Pred. No.	3.7e+04				
Matches	12	Conservative	0	Mismatches	0	Indels	0
Gaps	0						

Qy 1 GCCAGCGGAGTC 12
 |||||||
 Db 21 GCCAGCGGAGTC 10

<!--EndFragment-->